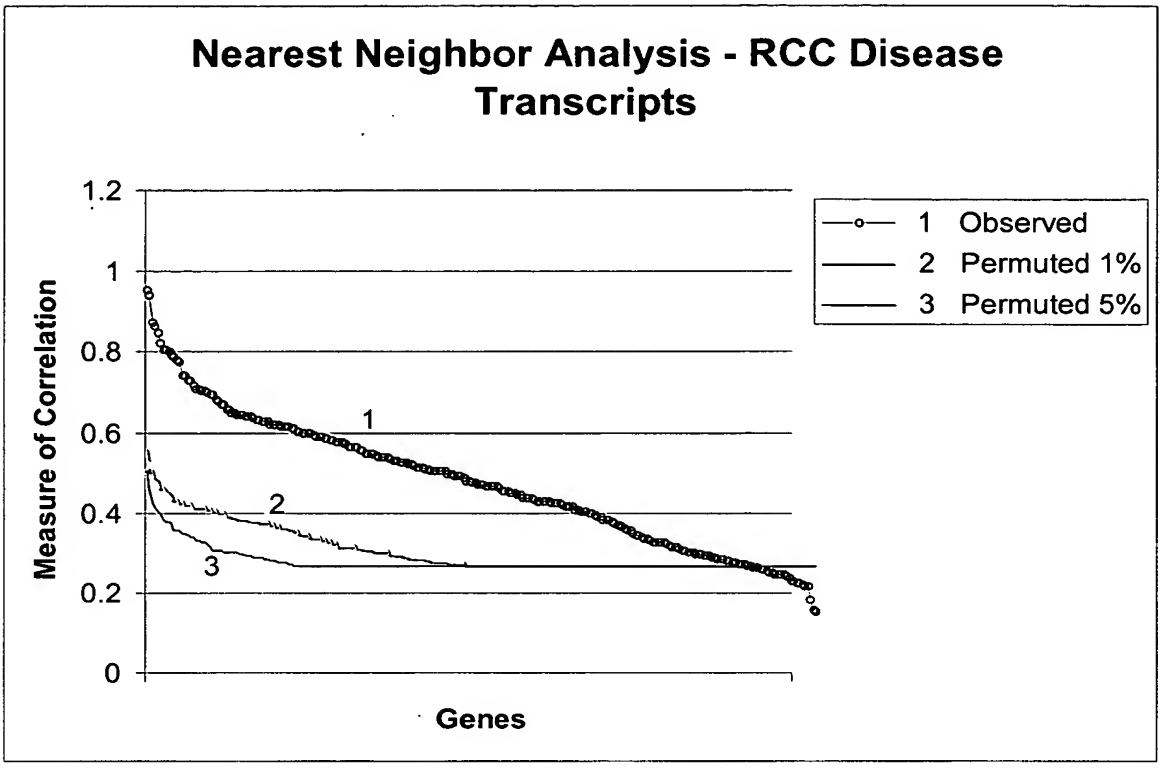

Nearest Neighbor Analysis - RCC Disease Transcripts
Measure of Correlation
1.2
1
0.8
0.6
0.4
0.2
0
Genes
1
2
3
1 Observed
2 Permuted 1%
3 Permuted 5%

FIG. 1

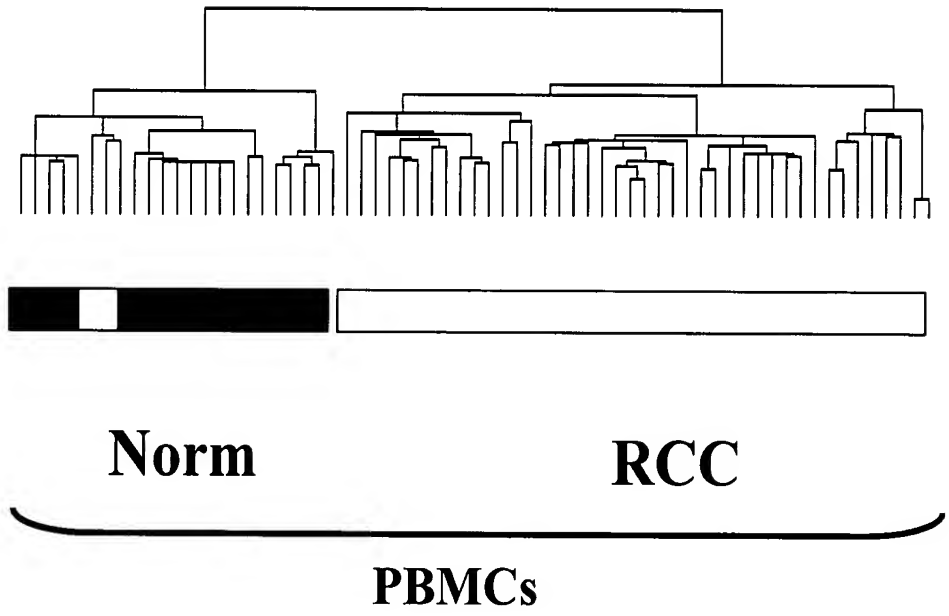


FIG. 2

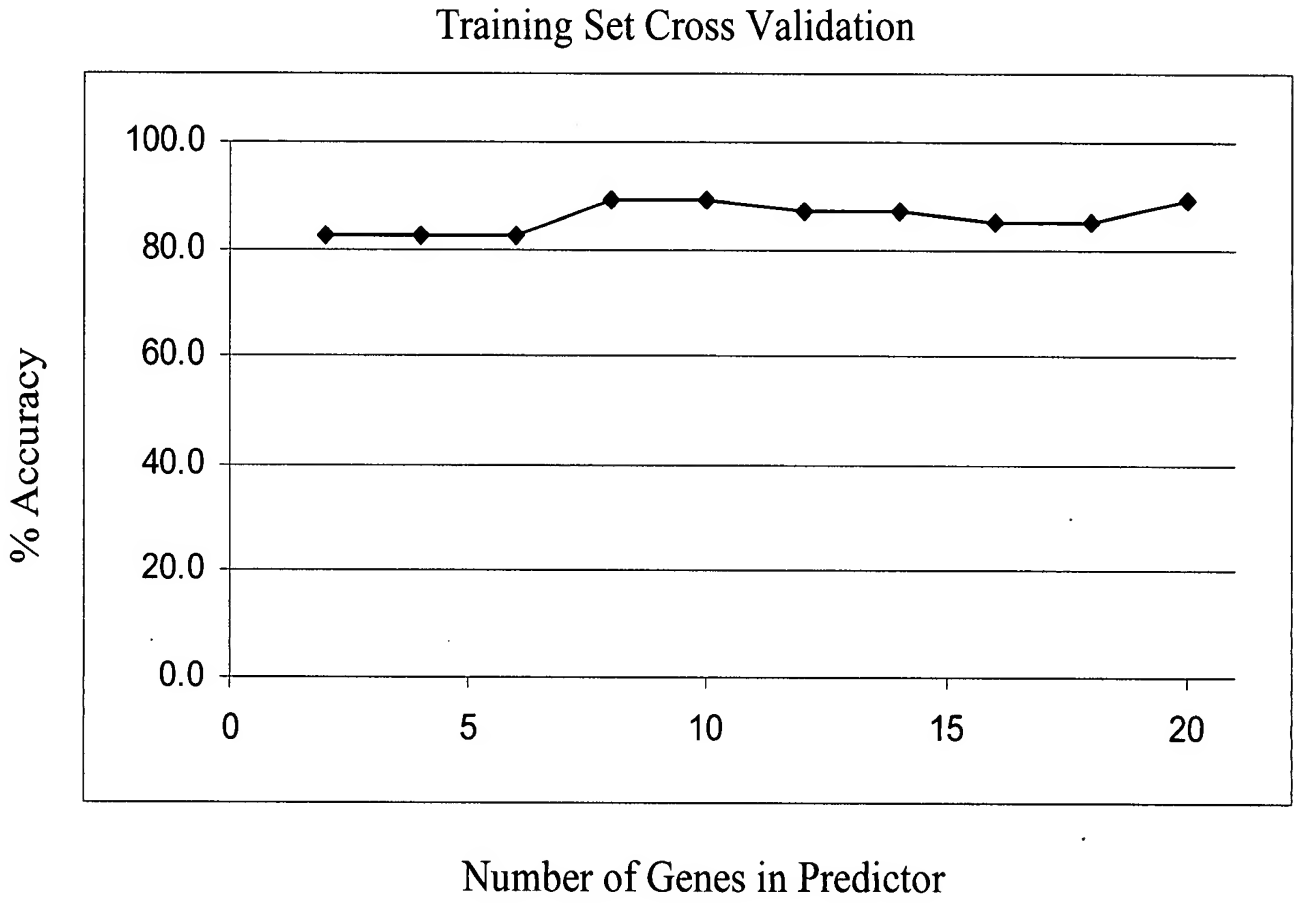

Training Set Cross Validation
100.0
80.0
60.0
40.0
20.0
0.0
% Accuracy
0
5
10
15
20
Number of Genes in Predictor

FIG. 3

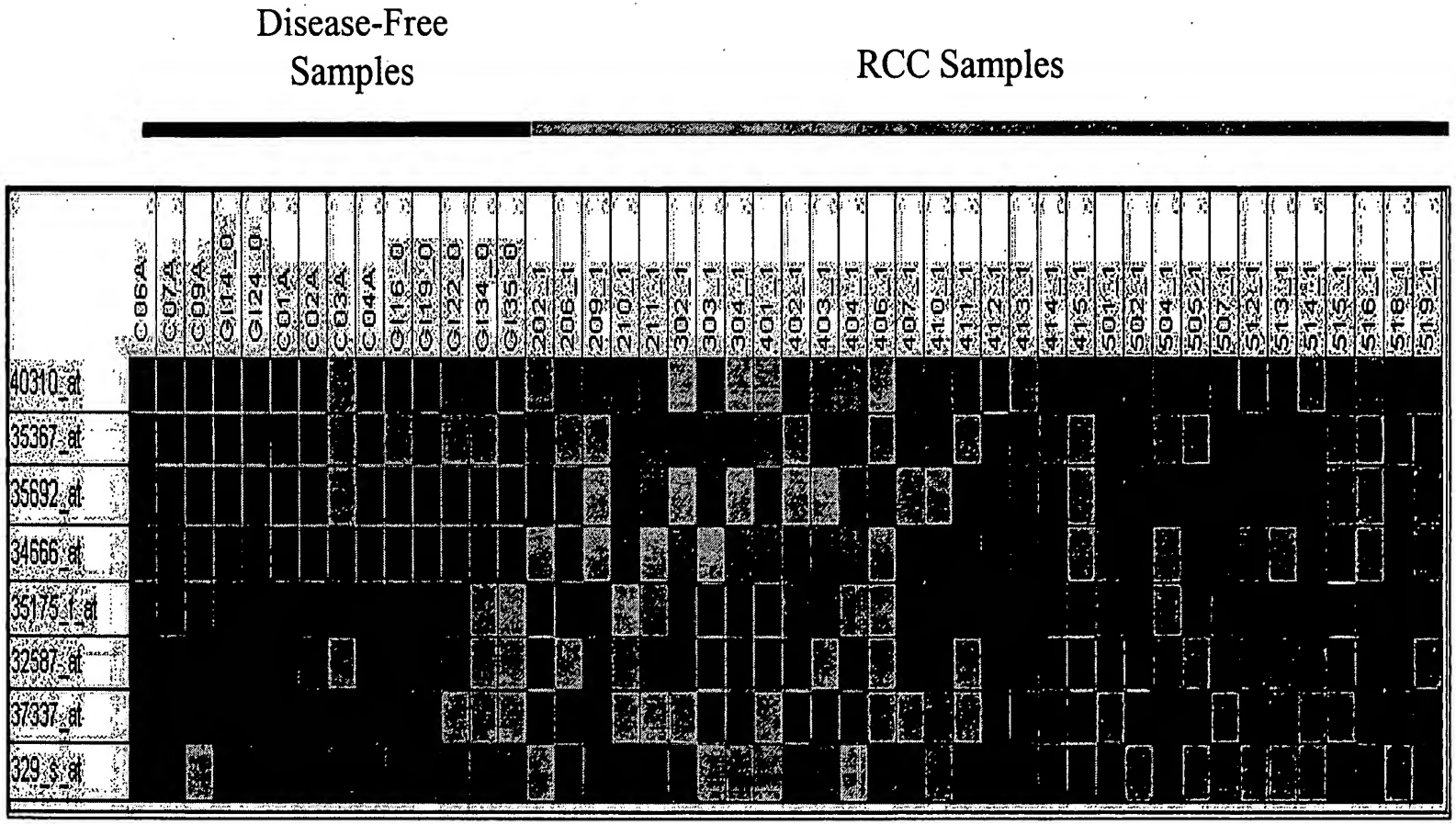
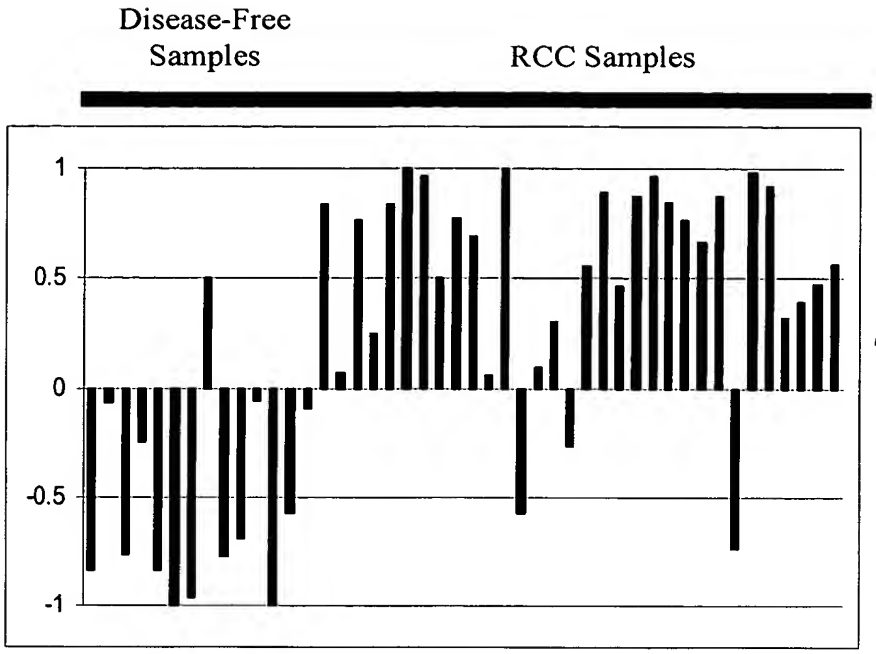


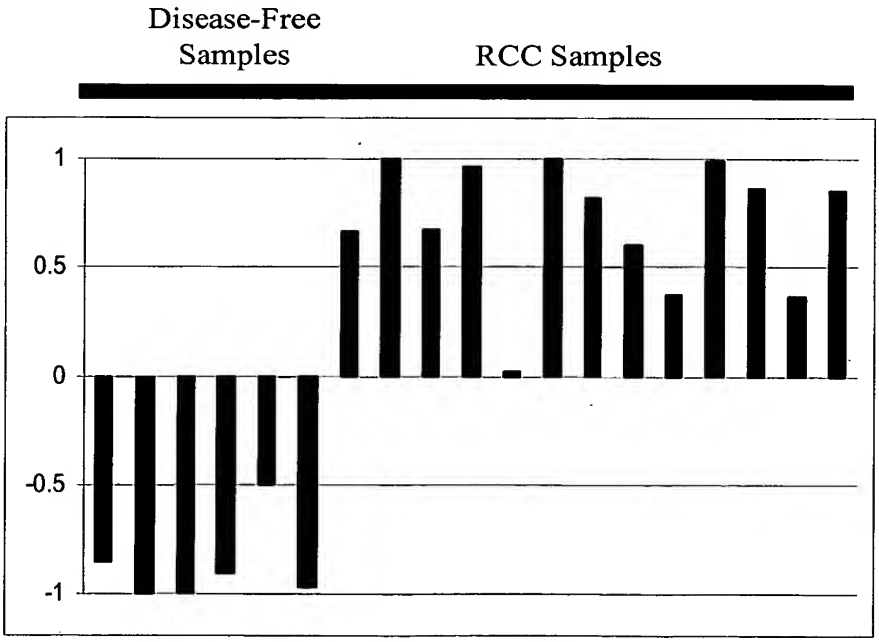
FIG. 4

BEST AVAILABLE COPY



Training Set
89%

FIG. 5A



Test Set
100%

FIG. 5B